

## SEQUENCE LISTING

<110> The Corporation of the Trustees of the Order of the Sisters of Mercy in Queensland

<120> Therapeutic and Diagnostic Agents

<130> 12373860/EJH

<150> AU 2002952993

<151> 2002-11-29

<160> 28

<170> PatentIn version 3.1

<210> 1

<211> 1151

<212> DNA

<213> Homo sapiens

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<212> PRT

<213> Homo sapiens

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35 40 45

His Arg Thr Leu Asn Lys Phe Trp Cys Arg Pro Pro Gln Ile Leu Arg  
50 55 60

Cys Asp Lys Ile Val Glu Thr Lys Gly Ser Ala Gly Lys Arg Asn Gly  
65 70 75 80

Arg Val Ser Ile Arg Asp Ser Pro Ala Asn Leu Ser Phe Thr Val Thr  
85 90 95

Leu Glu Asn Leu Thr Glu Glu Asp Ala Gly Thr Tyr Trp Cys Gly Val  
 100 105 110

Asp Thr Pro Trp Leu Arg Asp Phe His Asp Pro Ile Val Glu Val Glu  
 115 120 125

Val Ser Val Phe Pro Ala Gly Thr Thr Thr Ala Ser Ser Pro Gln Ser  
 130 135 140

Ser Met Gly Thr Ser Gly Pro Pro Thr Lys Leu Pro Val His Thr Trp  
 145 150 155 160

Pro Ser Val Thr Arg Lys Asp Ser Pro Glu Pro Ser Pro His Pro Gly  
 165 170 175

Ser Leu Phe Ser Asn Val Arg Phe Leu Leu Leu Val Leu Leu Glu Leu  
 180 185 190

Pro Leu Leu Leu Ser Met Leu Gly Ala Val Leu Trp Val Asn Arg Pro  
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<210> 3

<211> 1560

<212> DNA

<213> Homo sapiens

<400> 3

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<210> 4

<211> 301

<212> PRT

<213> Homo sapiens

<400> 4

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20 25 30

Val Gln Cys Pro Tyr Glu Lys Glu His Arg Thr Leu Asn Lys Tyr Trp  
35 40 45

Cys Arg Pro Pro Gln Ile Phe Leu Cys Asp Lys Ile Val Glu Thr Lys  
50 55 60

Gly Ser Ala Gly Lys Arg Asn Gly Arg Val Ser Ile Arg Asp Ser Pro  
65 70 75 80

Ala Asn Leu Ser Phe Thr Val Thr Leu Glu Asn Leu Thr Glu Glu Asp  
85 90 95

Ala Gly Thr Tyr Trp Cys Gly Val Asp Thr Pro Trp Leu Arg Asp Phe  
100 105 110

His Asp Pro Val Val Glu Val Glu Val Ser Val Phe Pro Ala Ser Thr  
115 120 125

Ser Met Thr Pro Ala Ser Ile Thr Ala Ala Lys Thr Ser Thr Ile Thr  
130 135 140

Thr Ala Phe Pro Pro Val Ser Ser Thr Thr Leu Phe Ala Val Gly Ala  
145 150 155 160

Thr His Ser Ala Ser Ile Gln Glu Glu Thr Glu Glu Val Val Asn Ser  
165 170 175

Gln Leu Pro Leu Leu Leu Ser Leu Leu Ala Leu Leu Leu Leu Leu Leu  
180 185 190

Val Gly Ala Ser Leu Leu Ala Trp Arg Met Phe Gln Lys Trp Ile Lys  
195 200 205

Trp Ile Lys Ala Gly Asp His Ser Glu Leu Ser Gln Asn Pro Lys Gln  
210 215 220

Ala Ala Thr Gln Ser Glu Leu His Tyr Ala Asn Leu Glu Leu Leu Met  
225 230 235 240

Trp Pro Leu Gln Glu Lys Pro Ala Pro Pro Arg Glu Val Glu Val Glu  
245 250 255

Tyr Ser Thr Val Ala Ser Pro Arg Glu Glu Leu His Tyr Ala Ser Val  
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Val Phe Asp Ser Asn Thr Asn Arg Ile Ala Ala Gln Arg Pro Arg Glu  
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Glu Glu Pro Asp Ser Asp Tyr Ser Val Ile Arg Lys Thr  
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<210> 5

<211> 674

<212> DNA

<213> Homo sapiens

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<210> 6

<211> 205

<212> PRT

<213> Homo sapiens

<400> 6

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Thr Val Trp Cys Gln Tyr Glu Ser Met Tyr Lys Gly Tyr Asn Lys Tyr  
35 40 45

Trp Cys Arg Gly Gln Tyr Asp Thr Ser Cys Glu Ser Ile Val Glu Thr  
50 55 60

Lys Gly Glu Glu Lys Val Glu Arg Asn Gly Arg Val Ser Ile Arg Asp  
65 70 75 80

His Pro Glu Ala Leu Ala Phe Thr Val Thr Met Gln Asn Leu Asn Glu  
85 90 95

Asp Asp Ala Gly Ser Tyr Trp Cys Lys Ile Gln Thr Val Trp Val Leu  
100 105 110

Asp Ser Trp Ser Arg Asp Pro Ser Asp Leu Val Arg Val Tyr Val Ser  
115 120 125

Pro Ala Ile Thr Thr Pro Arg Arg Thr Thr His Pro Ala Thr Pro Pro  
130 135 140

Ile Phe Leu Val Val Asn Pro Gly Arg Asn Leu Ser Thr Arg Glu Val  
145 150 155 160

Leu Thr Gln Asn Ser Gly Phe Arg Leu Ser Ser Pro His Phe Leu Leu  
165 170 175

Val Val Leu Leu Lys Leu Pro Leu Leu Leu Ser Met Leu Gly Ala Val  
180 185 190

Phe Trp Val Asn Arg Pro Gln Trp Ala Pro Pro Gly Arg  
195 200 205

<210> 7

<211> 510

<212> DNA

<213> Homo sapiens

<400> 7

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<210> 8

<211> 174

<212> PRT

<213> Homo sapiens

<400> 8

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20 25 30



Gly Ser Glu Gln Gly Ser Leu Thr Val Gln Cys Ala Tyr Gly Ser Gly  
35 40 45

Trp Glu Thr Tyr Leu Lys Trp Arg Cys Gln Gly Ala Asp Trp Asn Tyr  
50 55 60

Cys Asn Ile Leu Val Lys Thr Asn Gly Ser Glu Gln Glu Val Lys Lys  
65 70 75 80

Asn Arg Val Ser Ile Arg Asp Asn Gln Lys Asn His Val Phe Thr Val  
85 90 95

Thr Met Glu Asn Leu Lys Arg Asp Asp Ala Asp Ser Tyr Trp Cys Gly  
100 105 110

Thr Glu Arg Pro Gly Ile Asp Leu Gly Val Lys Val Gln Val Thr Ile  
115 120 125

Asn Pro Ala Gln Cys Leu Ser Leu Leu Pro Thr Asp Asp Arg Val Met  
130 135 140

Val Pro Val Ser Ala His Arg Pro Lys Gly Pro Pro Ser Leu Val Thr  
145 150 155 160

Arg Asp Pro Asn Pro Cys Gln Cys Leu Leu Gly Thr Ser Leu  
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<210> 9

<211> 1026

<212> DNA

<213> Homo sapiens

<400> 9

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<210> 10

<211> 193

<212> PRT

<213> Homo sapiens

<400> 10

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20 25 30

Val Gln Cys Arg Tyr Glu Asp Lys Tyr Lys Thr Phe Asn Lys Tyr Trp  
35 40 45

Cys Arg Gln Pro Cys Leu Pro Ile Trp His Glu Met Val Glu Thr Gly  
50 55 60

Gly Ser Glu Gly Val Val Arg Ser Asp Gln Val Ile Ile Thr Asp His  
65 70 75 80

Pro Gly Asp Leu Thr Phe Thr Val Thr Leu Glu Asn Leu Thr Ala Asp  
85 90 95

Asp Ala Gly Lys Tyr Arg Cys Gly Ile Ala Thr Ile Leu Gln Glu Asp  
100 105 110

Gly Leu Ser Gly Phe Leu Pro Asp Pro Phe Phe Gln Val Gln Val Leu  
115 120 125

Val Ser Ser Ala Ser Ser Thr Glu Asn Ser Val Lys Thr Pro Ala Ser  
130 135 140

Pro Thr Arg Pro Ser Gln Cys Gln Gly Ser Leu Pro Ser Ser Thr Cys  
145 150 155 160

Phe Leu Leu Leu Pro Leu Leu Lys Val Pro Leu Leu Leu Ser Ile Leu  
165 170 175

Gly Ala Ile Leu Trp Val Asn Arg Pro Trp Arg Thr Pro Trp Thr Glu  
180 185 190

Ser

<210> 11

<211> 1352

<212> DNA

<213> Homo sapiens

<400> 11

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<210> 12

<211> 158

<212> PRT

<213> Homo sapiens

<400> 12

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Met Trp Leu Pro Pro Ala Leu Leu Leu Ser Leu Ser Gly Cys Phe
1           5           10           15

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Thr Val Gln Cys His Tyr Lys Gln Gly Trp Glu Thr Tyr Ile Lys Trp  
35 40 45

Trp Cys Arg Gly Val Arg Trp Asp Thr Cys Lys Ile Leu Ile Glu Thr  
50 55 60

Arg Gly Ser Glu Gln Gly Glu Lys Ser Asp Arg Val Ser Ile Lys Asp  
65 70 75 80

Asn Gln Lys Asp Arg Thr Phe Thr Val Thr Met Glu Gly Leu Arg Arg  
85 90 95

Asp Asp Ala Asp Val Tyr Trp Cys Gly Ile Glu Arg Arg Gly Pro Asp  
100 105 110

Leu Gly Thr Gln Val Lys Val Ile Val Asp Pro Glu Gly Ala Ala Ser  
115 120 125

Thr Thr Ala Ser Ser Pro Thr Asn Ser Asn Met Ala Val Phe Ile Gly  
130 135 140

Ser His Lys Arg Asn His Tyr Met Leu Leu Gly Thr Ser Leu  
145 150 155

<210> 13

<211> 812

<212> DNA

<213> Homo sapiens

<400> 13

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<210> 14

<211> 287

<212> PRT

<213> Homo sapiens

<400> 14

Met Pro Leu Leu Thr Leu Tyr Leu Leu Leu Phe Trp Leu Ser Gly Tyr  
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Ser Ile Val Thr Gln Ile Thr Gly Pro Thr Thr Val Asn Gly Leu Glu  
20 25 30

Arg Gly Ser Leu Thr Val Gln Cys Val Tyr Arg Ser Gly Trp Glu Thr  
35 40 45

Tyr Leu Lys Trp Trp Cys Arg Gly Ala Ile Trp Arg Asp Cys Lys Ile  
50 55 60

Leu Val Lys Thr Ser Gly Ser Glu Gln Glu Val Lys Arg Asp Arg Val  
65 70 75 80

Ser Ile Lys Asp Asn Gln Lys Asn Arg Thr Phe Thr Val Thr Met Glu  
85 90 95

Asp Leu Met Lys Thr Asp Ala Asp Thr Tyr Trp Cys Gly Ile Glu Lys  
100 105 110

Thr Gly Asn Asp Leu Gly Val Thr Val Gln Val Thr Ile Asp Pro Ala  
115 120 125

Pro Val Thr Gln Glu Glu Thr Ser Ser Ser Pro Thr Leu Thr Gly His  
130 135 140

His Leu Asp Asn Arg His Lys Leu Leu Lys Leu Ser Val Leu Leu Pro  
145 150 155 160

Leu Ile Phe Thr Ile Leu Leu Leu Leu Leu Val Ala Ala Ser Leu Leu  
165 170 175

Ala Trp Arg Met Met Lys Tyr Gln Gln Lys Gly Glu Arg Thr Trp Val  
180 185 190

Leu Gln Pro Leu Glu Gly Asp Leu Cys Tyr Ala Asp Leu Thr Leu Gln  
195 200 205

Leu Ala Gly Thr Ser Pro Gln Lys Ala Thr Thr Lys Leu Ser Ser Ala  
210 215 220

Gln Val Asp Gln Val Glu Val Glu Tyr Val Thr Met Ala Ser Leu Pro  
225 230 235 240

Lys Glu Asp Ile Ser Tyr Ala Ser Leu Thr Leu Gly Ala Glu Asp Gln  
245 250 255

Glu Pro Thr Tyr Cys Asn Met Gly His Leu Ser Ser His Leu Pro Gly  
260 265 270

Arg Gly Pro Glu Glu Pro Thr Glu Tyr Ser Thr Ile Ser Arg Pro  
275 280 285

<210> 15

<211> 2389

<212> DNA

<213> mouse

<400> 15

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<210> 16

<211> 287

<212> PRT

<213> mouse

<400> 16

Met Arg Pro Leu Val Leu Leu Trp Gly Cys Leu Val Leu Pro Gly Tyr  
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Glu Ala Leu Lys Gly Pro Lys Glu Ile Ser Gly Phe Glu Gly Asp Thr  
20 25 30

Val Ser Leu Arg Cys Thr Tyr Val Glu Lys Met Lys Glu His Arg Lys  
35 40 45

Tyr Trp Cys Arg Gln Gly Gly Ile Leu Val Ser Arg Cys Gly Asp Ile  
50 55 60

Val Tyr Ala Asn Gln Asp Gln Glu Val Thr Arg Gly Arg Met Ser Ile  
65 70 75 80

Arg Asp Ser Pro Gln Glu Leu Ser Met Thr Val Ile Met Arg Asp Leu  
85 90 95

Thr Leu Lys Asp Ser Gly Lys Tyr Trp Cys Gly Ile Asp Arg Leu Gly  
100 105 110

Arg Asp Glu Ser Phe Glu Val Thr Leu Ile Val Phe Pro Gly Ser Ser  
115 120 125

Arg Pro Val Val Trp Leu Pro Leu Thr Thr Pro Gln Asp Ser Arg Ala  
130 135 140

Val Ala Ser Ser Val Ser Lys Pro Ser Val Ser Ile Pro Met Val Arg  
145 150 155 160

Met Met Ala Pro Val Leu Ile Leu Leu Ser Leu Leu Leu Ala Ala Gly  
165 170 175

Leu Ile Ala Phe Gly Ser His Met Leu Arg Trp Arg Lys Lys Ala Trp  
180 185 190

Leu Ala Thr Glu Thr Gln Lys Asn Glu Lys Val Tyr Leu Glu Thr Ser  
195 200 205

Leu Pro Gly Asn Gly Trp Thr Thr Glu Asp Ser Thr Ile Asp Leu Ala  
210 215 220

Val Thr Pro Glu Cys Leu Arg Asn Leu Asn Pro Ser Ala Val Pro Ser  
225 230 235 240

Pro Glu Thr Gln Asn Leu Ser Gln Ser Thr Glu Glu Glu Glu Ala Ala  
245 250 255

Arg Ser Leu Asp Asp Asp Lys Glu Asp Val Met Ala Pro Pro Pro Leu  
260 265 270

Gln Met Ser Ala Glu Glu Leu Ala Phe Ser Glu Phe Ile Ser Val  
275 280 285

<210> 17

<211> 1111

<212> DNA

<213> mouse

<400> 17

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caggattctc acgccaatgg agattctctt catcaacctc aggaccagaa agcagagtac      900
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<210> 18

<211> 314

<212> PRT

<213> mouse

<400> 18

Met Thr Gln Leu Ala Ser Ala Val Trp Leu Pro Thr Leu Leu Leu Leu  
1 5 10 15

Leu Leu Leu Phe Trp Leu Pro Gly Cys Val Pro Leu His Gly Pro Ser  
20 25 30

Thr Met Thr Gly Ser Val Gly Gln Ser Leu Ser Val Ser Cys Gln Tyr  
35 40 45

Glu Glu Lys Phe Lys Thr Lys Asp Lys Tyr Trp Cys Arg Gly Ser Leu  
50 55 60

Lys Val Leu Cys Lys Asp Ile Val Lys Thr Ser Ser Ser Glu Glu Ala  
65 70 75 80

Arg Ser Gly Arg Val Thr Ile Arg Asp His Pro Asp Asn Leu Thr Phe  
85 90 95

Thr Val Thr Tyr Glu Ser Leu Thr Leu Glu Asp Ala Asp Thr Tyr Met  
100 105 110

Cys Ala Val Asp Ile Ser Leu Phe Asp Gly Ser Leu Gly Phe Asp Lys  
115 120 125

Tyr Phe Lys Ile Glu Leu Ser Val Val Pro Ser Glu Asp Pro Gly Pro  
130 135 140

Thr Leu Glu Thr Pro Val Val Ser Thr Ser Leu Pro Thr Lys Gly Pro  
145 150 155 160

Ala Leu Gly Ser Asn Thr Glu Asp Arg Arg Glu His Asp Tyr Ser Gln  
165 170 175

Gly Leu Arg Leu Pro Ala Leu Leu Ser Val Leu Ala Leu Leu Leu Phe  
180 185 190

Leu Leu Val Gly Thr Ser Leu Leu Ala Trp Arg Met Phe Gln Lys Arg  
195 200 205

Leu Val Lys Ala Asp Arg His Pro Glu Leu Ser Gln Asn Leu Arg Gln  
210 215 220

Ala Ser Glu Gln Asn Glu Cys Gln Tyr Val Asn Leu Gln Leu His Thr  
225 230 235 240

Trp Ser Leu Arg Glu Glu Pro Val Leu Pro Ser Gln Val Glu Val Val  
245 250 255

Glu Tyr Ser Thr Leu Ala Leu Pro Gln Glu Glu Leu His Tyr Ser Ser  
260 265 270

Val Ala Phe Asn Ser Gln Arg Gln Asp Ser His Ala Asn Gly Asp Ser  
275 280 285

Leu His Gln Pro Gln Asp Gln Lys Ala Glu Tyr Ser Glu Ile Gln Lys  
290 295 300

Pro Arg Lys Gly Leu Ser Asp Leu Tyr Leu  
305 310

<210> 19

<211> 711

<212> DNA

<213> mouse

<400> 19  
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tggacggcag tacagacatg gtgtcttctg acttgcagaa gaggacttga agccagtcta 540  
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tttaccttcg tgctaacact gactcctcct agttcccagg aagcacacag cacaccgtca 660  
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<210> 20

<211> 236

<212> PRT

<213> mouse

<400> 20

Met Trp Leu Ser Pro Ala Leu Leu Leu Leu Ser Phe Pro Gly Cys Leu  
1 5 10 15

Ser Ile Gln Gly Pro Ala Leu Val Arg Gly Pro Glu Gln Gly Ser Val  
20 25 30

Thr Val Gln Cys Arg Tyr Ser Ser Arg Trp Gln Thr Asn Lys Lys Trp  
35 40 45

Trp Cys Arg Gly Ala Ser Trp Ser Thr Cys Arg Val Leu Ile Arg Ser  
50 55 60

Thr Gly Ser Glu Lys Glu Thr Lys Ser Gly Arg Leu Ser Ile Arg Asp  
65 70 75 80

Asn Gln Lys Asn His Ser Phe Gln Val Thr Met Glu Met Leu Arg Gln  
85 90 95

Asn Asp Thr Asp Thr Tyr Trp Cys Gly Ile Glu Lys Phe Gly Thr Asp  
100 105 110

Arg Gly Thr Arg Val Lys Val Asn Val Tyr Phe Gly His Met Gln Thr  
115 120 125

Phe Phe Ser Ser Ala Ala Thr Leu Thr Pro Glu Arg Ala Ala Glu Met  
130 135 140

Trp Val Lys Ile Pro Cys Arg Leu Leu Ile Asn Phe Pro Gly Pro Leu  
145 150 155 160

Trp Thr Ala Val Gln Thr Trp Cys Leu Leu Thr Cys Arg Arg Gly Leu  
165 170 175

Glu Ala Ser Leu Val Gly Ala Phe Val Gly Gly Leu Met Gln Val Pro  
180 185 190

Ser Cys Ser Leu Ala Val Ala Ile Phe Thr Phe Val Leu Thr Leu Thr  
195 200 205

Pro Pro Ser Ser Gln Glu Ala His Ser Thr Pro Ser Ser His Ser Ala  
210 215 220

Pro Val Ala Ser Lys Glu Glu Met Asn Arg Leu Phe  
225 230 235

<210> 21

<211> 819

<212> DNA

<213> mouse

<400> 21

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ttttctggag acagtaccca gtgaggcagg aggatgaggc tatgtgcagg tctgctcctt 180  
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<210> 22

<211> 181

<212> PRT

<213> mouse

<400> 22

Met Arg Leu Cys Ala Gly Leu Leu Leu Leu Cys Phe Gln Gly Cys Leu  
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Ser Leu Thr Gly Pro Gly Ser Val Ser Gly Tyr Val Gly Gly Ser Leu  
20 25 30

Arg Val Gln Cys Gln Tyr Ser Pro Ser Tyr Lys Gly Tyr Met Lys Tyr  
35 40 45

Trp Cys Arg Gly Pro His Asp Thr Thr Cys Lys Thr Ile Val Glu Thr  
50 55 60

Asp Gly Ser Glu Lys Glu Lys Arg Ser Gly Pro Val Ser Ile Arg Asp  
65 70 75 80

His Ala Ala Asn Ser Thr Ile Thr Val Ile Met Glu Asp Leu Ser Glu  
85 90 95

Asp Asp Ala Gly Ser Tyr Trp Cys Lys Ile Gln Thr Ser Phe Ile Trp  
100 105 110

Asp Ser Trp Ser Arg Asp Pro Ser Val Ser Val Arg Val Asn Val Phe  
115 120 125



Pro Val Asn Ser Gly Gln Asn Leu Arg Ile Ser Thr Asn Val Met Phe  
130 135 140

Ile Phe Gln Leu Trp Ser Leu Leu Ser Ser Ile Gln Phe Gln Val Leu  
145 150 155 160

Val Phe Leu Lys Leu Pro Leu Phe Leu Ser Met Leu Cys Ala Ile Phe  
165 170 175

Trp Val Asn Arg Leu  
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<210> 23

<211> 2487

<212> DNA

<213> mouse

<400> 23

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gttctctgct ctactcctat tcttctctccc aggctgctgc acggctcagg attcagtcac 180  
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2487

<210> 24

<211> 221

<212> PRT

<213> mouse

<400> 24

Met Trp Gln Phe Ser Ala Leu Leu Leu Phe Phe Leu Pro Gly Cys Cys  
1 5 10 15

Thr Ala Gln Asp Ser Val Thr Gly Pro Glu Glu Val Ser Gly Gln Glu  
20 25 30

Gln Gly Ser Leu Thr Val Gln Cys Arg Tyr Ser Ser Tyr Trp Lys Gly  
35 40 45

Tyr Lys Lys Tyr Trp Cys Arg Gly Val Pro Gln Arg Ser Cys Asp Ile  
50 55 60

Leu Val Glu Thr Asp Lys Ser Glu Gln Leu Val Lys Lys Asn Arg Val  
65 70 75 80

Ser Ile Arg Asp Asn Gln Arg Asp Phe Ile Phe Thr Val Thr Met Glu  
85 90 95

Asp Leu Arg Met Ser Asp Ala Gly Ile Tyr Trp Cys Gly Ile Thr Lys  
100 105 110

Gly Gly Pro Asp Pro Met Phe Lys Val Asn Val Asn Ile Asp Gln Ala  
115 120 125

Pro Lys Ser Ser Met Met Thr Thr Thr Ala Thr Val Leu Lys Ser Ile  
130 135 140

Gln Pro Ser Ala Glu Asn Thr Gly Lys Glu Gln Val Thr Gln Ser Lys  
145 150 155 160

Glu Val Thr Gln Ser Arg Pro His Thr Arg Ser Leu Leu Ser Ser Ile  
 165 170 175

Tyr Phe Leu Leu Met Val Phe Val Glu Leu Pro Leu Leu Leu Ser Met  
 180 185 190

Leu Ser Ala Val Leu Trp Val Thr Arg Pro Gln Arg Cys Phe Gly Arg  
 195 200 205

Gly Glu Asn Asp Leu Val Lys Thr His Ser Pro Val Ala  
 210 215 220

<210> 25

<211> 1307

<212> DNA

<213> mouse

<400> 25

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tgagaagatg catttgtcat tgctgggtccc ctttctcttc tggatcacag gctgctgcac	180
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tcctcagaga tcatgtaaga ctcttggtga aaccgatgca tcagagcagc tgggtgaagaa	360
gaaccgtgtg tccatcaggg acaaccagag agacttcac ttcacagtga ccatggagga	420
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tgggtgaagat ggcgtcggtg atgggtttct ggatctcagt gtgctcctcc cagtcacttc	660
tgagtcctg ttgcttctcc tggttggtggc ctgctctttt gcttgaggga tggtgaggag	720
acagaagaaa gacctgtccc tgaagcagcc cagaacctcc cctggctcct cttggaaaaa	780
gggtcctccc atgtcctcct ctggcaagga ccaccaagag gaagtggaat atgtcaccat	840
ggctcccttt ccagggagg aggtttcata tgccgctctg actttggccg gcttggttca	900

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ggagcctact tatggcaata ctggctgccc catcacccat gttcccagga caggccttga      960
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ttggtctctg aaggcggctt ggagcatgga tctttacatc tgcctctgta cctgcttcct      1080
taccgggccc agctggtgac tggaactctg tccatccgtc tctcatggcc atcagctcta      1140
ccttgcttga gcttggagtt caacctcagg gggttccagg gaattaaggc tccttccaca      1200
tccccactta tagccaatgt accttggaag gtaccaggca ggctgcttca gggatgctgt      1260
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<210> 26

<211> 296

<212> PRT

<213> mouse

<400> 26

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Met His Leu Ser Leu Leu Val Pro Phe Leu Phe Trp Ile Thr Gly Cys
1              5              10              15

```

```

Cys Thr Ala Glu Asp Pro Val Thr Gly Pro Glu Glu Val Ser Gly Gln
                20              25              30

```

```

Glu Gln Gly Ser Leu Thr Val Gln Cys Arg Tyr Thr Ser Gly Trp Lys
35              40              45

```

```

Asp Tyr Lys Lys Tyr Trp Cys Gln Gly Val Pro Gln Arg Ser Cys Lys
50              55              60

```

```

Thr Leu Val Glu Thr Asp Ala Ser Glu Gln Leu Val Lys Lys Asn Arg
65              70              75              80

```

```

Val Ser Ile Arg Asp Asn Gln Arg Asp Phe Ile Phe Thr Val Thr Met
85              90              95

```

```

Glu Asp Leu Arg Met Ser Asp Ala Gly Ile Tyr Trp Cys Gly Ile Thr
100              105              110

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Lys Val Pro Thr Met Pro Pro Ile Thr Ser Thr Thr Thr Ile Phe Thr  
115 120 125

Val Thr Thr Thr Val Lys Glu Thr Ser Met Phe Pro Thr Leu Thr Ser  
130 135 140

Tyr Tyr Ser Asp Asn Gly His Gly Gly Gly Asp Ser Gly Gly Gly Glu  
145 150 155 160

Asp Gly Val Gly Asp Gly Phe Leu Asp Leu Ser Val Leu Leu Pro Val  
165 170 175

Ile Ser Ala Val Leu Leu Leu Leu Leu Leu Val Ala Ser Leu Phe Ala  
180 185 190

Trp Arg Met Val Arg Arg Gln Lys Lys Asp Leu Ser Leu Lys Gln Pro  
195 200 205

Arg Thr Ser Pro Gly Ser Ser Trp Lys Lys Gly Ser Ser Met Ser Ser  
210 215 220

Ser Gly Lys Asp His Gln Glu Glu Val Glu Tyr Val Thr Met Ala Pro  
225 230 235 240

Phe Pro Arg Glu Glu Val Ser Tyr Ala Ala Leu Thr Leu Ala Gly Leu  
245 250 255

Gly Gln Glu Pro Thr Tyr Gly Asn Thr Gly Cys Pro Ile Thr His Val  
260 265 270

Pro Arg Thr Gly Leu Glu Glu Glu Thr Thr Glu Tyr Ser Ser Ile Arg  
275 280 285

Arg Pro Leu Pro Ala Ala Met Pro  
290 295

<210> 27

<211> 114

<212> PRT

<213> mouse

<400> 27

Gly Cys Cys Thr Ala Gln Asp Pro Val Thr Gly Pro Glu Glu Val Ser  
1 5 10 15

Gly Gln Glu Gln Gly Ser Leu Thr Val Gln Cys Arg Tyr Asp Ser Gly  
20 25 30

Trp Lys Asp Tyr Lys Lys Tyr Trp Cys Arg Gly Ala Tyr Trp Lys Ser  
35 40 45

Cys Glu Ile Leu Val Glu Thr Asp Ala Ser Glu Gln Leu Val Lys Glu  
50 55 60

Asn Arg Val Ser Ile Arg Asp Asp Gln Thr Asp Phe Ile Phe Thr Val  
65 70 75 80

Thr Met Glu Asp Leu Arg Met Ser Asp Ala Asp Ile Tyr Trp Cys Gly  
85 90 95

Ile Thr Lys Ala Gly Thr Asp Pro Met Phe Lys Val Asn Val Asn Ile  
100 105 110

Asp Pro

<210> 28

<211> 295

<212> PRT

<213> homosapiens

<400> 28

Met Pro Leu Leu Thr Leu Tyr Leu Leu Leu Phe Trp Leu Ser Gly Tyr  
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Ser Ile Val Thr Gln Ile Thr Gly Pro Thr Thr Val Asn Gly Leu Glu  
20 25 30

Arg Gly Ser Leu Thr Val Gln Cys Val Tyr Arg Ser Gly Trp Glu Thr  
35 40 45

Tyr Leu Lys Trp Trp Cys Arg Gly Ala Ile Trp Arg Asp Cys Lys Ile  
50 55 60

Leu Val Lys Thr Ser Gly Ser Glu Gln Glu Val Lys Arg Asp Arg Val  
65 70 75 80

Ser Ile Lys Asp Asn Gln Lys Asn Arg Thr Phe Thr Val Thr Met Glu  
85 90 95

Asp Leu Met Lys Thr Asp Ala Asp Thr Tyr Trp Cys Gly Ile Glu Lys  
100 105 110

Thr Gly Asn Asp Leu Gly Val Thr Val Gln Val Thr Ile Asp Pro Ala  
115 120 125

Pro Val Thr Gln Glu Glu Thr Ser Ser Ser Pro Thr Leu Thr Gly His  
130 135 140

His Leu Asp Asn Arg His Lys Leu Leu Lys Leu Ser Val Leu Leu Pro  
145 150 155 160

Leu Ile Phe Thr Ile Leu Leu Leu Leu Leu Val Ala Ala Ser Leu Leu  
165 170 175

Ala Trp Arg Met Met Lys Tyr Gln Gln Lys Gly Glu Arg Thr Trp Val  
180 185 190

Leu Gln Pro Leu Glu Gly Asp Leu Cys Tyr Ala Asp Leu Thr Leu Gln  
195 200 205

Leu Ala Gly Thr Ser Pro Gln Lys Ala Thr Thr Lys Leu Ser Ser Ala  
210 215 220

Gln Val Asp Gln Val Glu Val Glu Tyr Val Ala Ala Gly Met Ser Pro  
225 230 235 240

Glu Gln Thr Met Ala Ser Leu Pro Lys Glu Asp Ile Ser Tyr Ala Ser  
245 250 255



Leu Thr Leu Gly Ala Glu Asp Gln Glu Pro Thr Tyr Cys Asn Met Gly  
260 265 270

His Leu Ser Ser His Leu Pro Gly Arg Gly Pro Glu Glu Pro Thr Glu  
275 280 285

Tyr Ser Thr Ile Ser Arg Pro  
290 295